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 United States Patent and Trademark Office
 Washington, D.C. 20231
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U.S. APPLICATION NO.	FIRST NAMED APPLICANT	ATTY. DOCKET NO.
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09/763994

FIRMAN'S

B X-12239

ELI LILLY AND COMPANY
 LILLY CORPORATE CENTER
 INDIANAPOLIS, IN 46285

RECEIVED

OCT 02 2001

ELI LILLY & COMPANY
 PATENT DIVISION

INTERNATIONAL APPLICATION NO.

PCT/US99/19436

I.A. FILING DATE	PRIORITY DATE
------------------	---------------

30 AUG 99

01 AUG 98

DATE MAILED: 20 SEP 2001

NOTIFICATION OF A DEFECTIVE RESPONSE

1. The request for an extension of time (37 CFR 1.136(a)) filed _____ is defective because the required fee is missing/insufficient. Extension of time fees are listed at 37 CFR 1.17(a)(1)-(a)(5).
2. Applicant's response filed _____ was received in the Office after the expiration of the period for response set in the Office notification mailed _____. This application will become abandoned unless applicant obtains an extension of time to reply to the last Office notification under 37 CFR 1.136(a).
3. Applicant's response filed JUN 08 2001 is hereby acknowledged. The following requirements set forth in the NOTIFICATION of MISSING REQUIREMENTS (Form PCT/DO/EO/905) mailed APR 02 2001 have not been completed.

- Translation of the international application into English.
 - which is defective for the reasons indicated on the attached Notice of Defective Translation.
- Processing fee (37 CFR 1.492(f)).
- Oath or Declaration of inventor(s).
 - not in compliance with 37 CFR 1.497(a) and (b) for the reasons indicated on the attached PCT/DO/EO/917.
- SurchARGE (37 CFR 1.492(c)).
- Sequence Listing.
 - not in compliance with 37 CFR 1.821-1.825 for the reasons indicated on the attached PCT/DO/EO/920.
- Additional claim fees.

Applicant is required to complete the response within a time limit of ONE MONTH from the date of this Notification or within the time remaining in the response set forth in the Notification of Missing Requirements (Form DO/EO/905), whichever is the longer. No extension of this time limit may be granted under 37 C.F.R. § 1.136, but the period for response set in the Notification of Missing Requirements (Form DO/EO/905) may be extended under 37 C.F.R. § 1.136(a).

Applicant is reminded that any communication to the United States Patent and Trademark Office must be mailed to the address given in the heading and include the U.S. application no. shown above. (37 CFR 1.5)

Enclosed: PCT/DO/EO/917 Notice of Defective Translation
 PCT/DO/EO/920

John L. Anderson

Telephone: 703-308-9116



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U.S. APPLICATION NO.	FIRST NAMED APPLICANT	ATTY. DOCKET NO.
09/763994	EDMONDS	B X-12239
		INTERNATIONAL APPLICATION NO.
		PCT/US99/19436
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30 AUG 99		01 SEP 98

DATE MAILED:

20 SEP 2001

**NOTIFICATION TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS
 CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE
 DISCLOSURES**

Applicant has submitted papers under 35 U.S.C. 371 to enter the national stage in the United States of America. The items indicated below, however, are missing. The period within which to correct the deficiency noted below and avoid abandonment is set forth in the accompanying Notification.

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 CFR 1.821-1.825 for the following reason(s):

- The application fails to comply with the requirements of 37 CFR 1.821-1.825.
- This application does not contain, a "Sequence Listing" as a separate part of the disclosure on paper copy or compact disc, as required by 37 CFR 1.821(c).
- A copy of the "Sequence Listing" in computer readable format has not been submitted as required by 37 CFR 1.821(e).
- A copy of the "Sequence Listing" in computer readable form has been submitted. The content of the computer readable form, however, does not comply with the requirements of 37 CFR 1.822 and/or 1.832, as indicated on the attached marked-up copy of the "Raw Sequence Listing."
- The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A substitute computer readable form must be submitted as required by 37 CFR 1.825(d).
- The paper copy or compact disc of the "Sequence Listing" is not the same as the computer readable form of the "Sequence Listing" as required by 37 CFR 1.821(e).
- Other: _____

APPLICANT MUST PROVIDE:

- An initial or substitute computer readable form (CRF) of the "Sequence Listing."
- An initial or substitute paper copy or compact disc of the "Sequence Listing," as well as an amendment directing its entry into the specification.
- A statement that the contents of the paper or compact disc and the computer readable form are the same and, where applicable, include no new matter, as required by 37 CFR 1.821(e), 1.821(f), 1.821(g), 1.825(b) or 1.825(d).

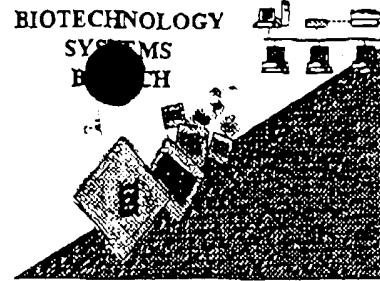
FOR QUESTIONS REGARDING COMPLIANCE WITH THESE REQUIREMENTS, PLEASE CALL:

- (703) 308-4216, for Rules interpretation,
- (703) 308-4212, for CRF submission help,
- (703) 287-0200, for PatentIn software help

John L. Anderson

Telephone: 703-308-9116

RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/263,994

Source: PCF09

Date Processed by STIC: 2/11/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

PCT09

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/763,994

DATE: 07/11/2001

TIME: 11:05:29

Input Set : A:\X-12239SeqList.app
 Output Set: N:\CRF3\07112001\I763994.raw

Does Not Comply
 Corrected Diskette Needed

3 <110> APPLICANT: Edmonds, Brian T.
 5 <120> TITLE OF INVENTION: HUMAN LATENT TRANSFORMING GROWTH FACTOR-BETA BINDING
 6 PROTEIN 3
 8 <130> FILE REFERENCE: X-12239
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/763,994
 C--> 11 <141> CURRENT FILING DATE: 2001-06-08
 13 <160> NUMBER OF SEQ ID NOS: 6
 15 <170> SOFTWARE: PatentIn Ver. 2.0
 17 <210> SEQ ID NO: 1
 18 <211> LENGTH: 3624
 19 <212> TYPE: DNA
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 24 gtgatctgca agcggacactg tctcaagggc cagtgtcggg acagttgtca gcagggtctcc 120
 25 aacatgacgc tcatacgaga gaacggccac agcacagaca cgctcacggg ctccggcttc 180
 26 cgcgtgggtgg tgtgcctct cccctgcatg aatggcgcc agtgcctctc gcgaaccagg 240
 27 tgccctgtgtc cccccgactt cactgggcgc ttctgccagg tgcccgcaagg aggagccgg 300
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 30 caggtgatcg ctgaccctcc tggggccggg gagggccctc ctgcccagca cgcagccctc 480
 31 ctgggtcccc taggccccggg acagatctca gcagaagtgc aggccccggc ccccggtgt 540
 32 aatgtgcgcg tccatcaccc gcccggaggcc tcagttccagg tgcacccgtat tgagagctcg 600
 33 aacgcccggaa gcccggggcc ctcccaagcac ctgctgccgc accccaagcc ctcgcaccccc 660
 34 cggccgccccca cccagaagtc cttggggccgc tgcttcagg acactctgcc caagcagccg 720
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 55 cccggctacc ggctcaaaggc ctccggccct cctgtgtgcg aagacatgca cgggtgcgg 1980
 56 gacccttgc tggcaatgc gagaacaaggc cccggagctt caagtgcatt 2040

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/763,994

07/11/2001
TIME: 11:05:29

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Output Set: N:\CRF3\07112001\I763994 raw

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 58 gcccaggca gcccctgtc gcctggctgg tgcgagaacc tcccgccgtc cttccgcgtc 2160
 59 acctgtgccc agggctacgc gcccggccccc gacggccgca gttgtcttggg tggacgacgag 2220
 60 tgtgaggctg gggacgtgt tgacaatggc atctgcagca acacgcccagg atctttccag 2280
 61 tgtcagtgc tctctggcta ccatactgtcc agggaccggg gccactgcga ggacattgtat 2340
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 64 gagtgccaggc aggaccccgag cctgtgcctt ccccatgggg cctgcaagaa ctttcaggc 2520
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 66 gaggtggagc agccccacca caagaaggag tgctacctga acttcgtatga cacagtgttc 2640
 67 tgcgacagcg tattggccac caacgtgacc cagcaggagt gtcgtgtc tctggggcc 2700
 68 ggctggggcg accactgcga aacttacccc tgcccagtct acagctcagc cgagttccac 2760
 69 agcctctgcc cagacggaaa gggctacacc caggacaaca acatcgtaa ctacggcatc 2820
 70 ccagcccacc gtgacatcgatcgatg ttgttcgggt cggagatgg caaggaggc 2880
 71 aagtgcgtga acacgcagcc tggctacgag tgctactgca agcagggtctt ctactacgac 2940
 72 gggAACCTGC tggaaatgcgt ggacgtggac gagggtccttgg accagttccaa ctggccggaa 3000
 73 ggagtgtgtg agaacacgcg cggcggctac cgctgtgcct gcacggcccc tgccgagttac 3060
 74 agtccccgcg acgcggcaggc cctgagcccg gaagagatgg agcgtggcccc ggagcggcgc 3120
 75 gacgtgtgt ggagccagcg cggagaggac ggcattgtcg ctggccccctt ggccggggcct 3180
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 77 cctgtccccgc cgcgcggcgc ggggtcccat tgccgcacat cgcagagcga gagcaattcc 3300
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 87 <212> TYPE: PRT
 88 <213> ORGANISM: Homo sapiens
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 91 Arg Gly Ala Gly Gly Gly Ala Leu Ala Arg Glu Arg Phe Lys Val
 92 1 5 10 15
 94 Val Phe Ala Pro Val Ile Cys Lys Arg Thr Cys Leu Lys Gly Gln Cys
 95 20 25 30
 97 Arg Asp Ser Cys Gln Gln Gly Ser Asn Met Thr Leu Ile Gly Glu Asn
 98 35 40 45
 100 Gly His Ser Thr Asp Thr Leu Thr Gly Ser Gly Phe Arg Val Val Val
 101 50 55 60
 103 Cys Pro Leu Pro Cys Met Asn Gly Gly Gln Cys Ser Ser Arg Asn Gln
 104 65 70 75 80
 106 Cys Leu Cys Pro Pro Asp Phe Thr Gly Arg Phe Cys Gln Val Pro Ala
 107 85 90 95
 109 Gly Gly Ala Gly Gly Gly Thr Gly Gly Ser Gly Pro Gly Leu Ser Arg
 110 100 105 110
 112 Thr Gly Ala Leu Ser Thr Gly Ala Leu Pro Pro Leu Ala Pro Glu Gly
 113 115 120 125
 115 Asp Ser Val Ala Ser Lys His Ala Ile Tyr Ala Val Gln Val Ile Ala

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/763,994

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Input Set : A:\X-12239SeqList.app
Output Set: N:\CRF3\07112001\I763994.raw

116	130	135	140
118	Asp Pro Pro Gly Pro Gly Glu Gly Pro Pro Ala Gln His Ala Ala Phe		
119	145	150	155
121	Leu Val Pro Leu Gly Pro Gly Gln Ile Ser Ala Glu Val Gln Ala Pro		160
122	165	170	175
124	Pro Pro Val Val Asn Val Arg Val His His Pro Pro Glu Ala Ser Val		
125	180	185	190
127	Gln Val His Arg Ile Glu Ser Ser Asn Ala Glu Ser Ala Ala Pro Ser		
128	195	200	205
130	Gln His Leu Leu Pro His Pro Lys Pro Ser His Pro Arg Pro Pro Thr		
131	210	215	220
133	Gln Lys Ser Leu Gly Arg Cys Phe Gln Asp Thr Leu Pro Lys Gln Pro		
134	225	230	235
136	Cys Gly Ser Asn Pro Leu Pro Gly Leu Thr Lys Gln Glu Asp Cys Cys		240
137	245	250	255
139	Gly Ser Ile Gly Thr Ala Trp Gly Gln Ser Lys Cys His Lys Cys Pro		
140	260	265	270
142	Gln Leu Gln Tyr Thr Gly Val Gln Lys Pro Gly Pro Val Arg Gly Glu		
143	275	280	285
145	Val Gly Ala Asp Cys Pro Gln Gly Tyr Lys Arg Leu Asn Ser Thr His		
146	290	295	300
148	Cys Gln Asp Ile Asn Glu Cys Ala Met Pro Gly Val Cys Arg His Gly		
149	305	310	315
151	Asp Cys Leu Asn Asn Pro Gly Ser Tyr Arg Cys Val Cys Pro Pro Gly		
152	325	330	335
154	His Ser Leu Gly Pro Ser Arg Thr Gln Cys Ile Ala Asp Lys Pro Glu		
155	340	345	350
157	Glu Lys Ser Leu Cys Phe Arg Leu Val Ser Pro Glu His Gln Cys Gln		
158	355	360	365
160	His Pro Leu Thr Thr Arg Leu Thr Arg Gln Leu Cys Cys Ser Val		
161	370	375	380
163	Gly Lys Ala Trp Gly Ala Arg Cys Gln Arg Cys Pro Thr Asp Gly Thr		
164	385	390	395
166	Ala Ala Phe Lys Glu Ile Cys Pro Ala Gly Lys Gly Tyr His Ile Leu		400
167	405	410	415
169	Thr Ser His Gln Thr Leu Thr Ile Gln Gly Glu Ser Asp Phe Ser Leu		
170	420	425	430
172	Phe Leu His Pro Asp Gly Pro Pro Lys Pro Gln Gln Leu Pro Glu Ser		
173	435	440	445
175	Pro Ser Gln Ala Pro Pro Pro Glu Asp Thr Glu Glu Glu Arg Gly Val		
176	450	455	460
178	Thr Thr Asp Ser Pro Val Ser Glu Glu Arg Ser Val Gln Gln Ser His		
179	465	470	475
181	Pro Thr Ala Thr Thr Pro Ala Arg Pro Tyr Pro Glu Leu Ile Ser		480
182	485	490	495
184	Arg Pro Ser Pro Pro Thr Met Arg Trp Phe Leu Pro Asp Leu Pro Pro		
185	500	505	510
187	Ser Arg Ser Ala Val Glu Ile Ala Pro Thr Gln Val Thr Glu Thr Asp		
188	515	520	525

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/763,994

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Input Set : A:\X-12239SeqList.app
Output Set: N:\CRF3\07112001\I763994.raw

190 Glu Cys Arg Leu Asn Gln Asn Ile Cys Gly His Gly Glu Cys Val Pro
191 530 535 540
193 Gly Pro Pro Asp Tyr Ser Cys His Cys Asn Pro Gly Tyr Arg Ser His
194 545 550 555 560
196 Pro Gln His Arg Tyr Cys Val Asp Val Asn Glu Cys Glu Ala Glu Pro
197 565 570 575
199 Cys Gly Pro Gly Arg Gly Ile Cys Met Asn Thr Gly Gly Ser Tyr Asn
200 580 585 590
202 Cys His Cys Asn Arg Gly Tyr Arg Leu His Val Gly Ala Gly Gly Arg
203 595 600 605
205 Ser Cys Val Asp Leu Asn Glu Cys Ala Lys Pro His Leu Cys Gly Asp
206 610 615 620
208 Gly Gly Phe Cys Ile Asn Phe Pro Gly His Tyr Lys Cys Asn Cys Tyr
209 625 630 635 640
211 Pro Gly Tyr Arg Leu Lys Ala Ser Arg Pro Pro Val Cys Glu Asp Ile
212 645 650 655
214 Asp Glu Cys Arg Asp Pro Ser Ser Cys Pro Asp Gly Lys Cys Glu Asn
215 660 665 670
217 Lys Pro Gly Ser Phe Lys Cys Ile Ala Cys Gln Pro Gly Tyr Arg Ser
218 675 680 685
220 Gln Gly Gly Ala Cys Arg Asp Val Asn Glu Cys Ala Glu Gly Ser
221 690 695 700
223 Pro Cys Ser Pro Gly Trp Cys Glu Asn Leu Pro Gly Ser Phe Arg Cys
224 705 710 715 720
226 Thr Cys Ala Gln Gly Tyr Ala Pro Ala Pro Asp Gly Arg Ser Cys Leu
227 725 730 735
229 Asp Val Asp Glu Cys Glu Ala Gly Asp Val Cys Asp Asn Gly Ile Cys
230 740 745 750
232 Ser Asn Thr Pro Gly Ser Phe Gln Cys Gln Cys Leu Ser Gly Tyr His
233 755 760 765
235 Leu Ser Arg Asp Arg Ser His Cys Glu Asp Ile Asp Glu Cys Asp Phe
236 770 775 780
238 Pro Ala Ala Cys Ile Gly Gly Asp Cys Ile Asn Thr Asn Gly Ser Tyr
239 785 790 795 800
241 Arg Cys Leu Cys Pro Gln Gly His Arg Leu Val Gly Gly Arg Lys Cys
242 805 810 815
244 Gln Asp Ile Asp Glu Cys Ser Gln Asp Pro Ser Leu Cys Leu Pro His
245 820 825 830
247 Gly Ala Cys Lys Asn Leu Gln Gly Ser Tyr Val Cys Val Cys Asp Glu
248 835 840 845
250 Gly Phe Thr Pro Thr Gln Asp Gln His Gly Cys Glu Glu Val Glu Gln
251 850 855 860
253 Pro His His Lys Lys Glu Cys Tyr Leu Asn Phe Asp Asp Thr Val Phe
254 865 870 875 880
256 Cys Asp Ser Val Leu Ala Thr Asn Val Thr Gln Gln Glu Cys Cys Cys
257 885 890 895
259 Ser Leu Gly Ala Gly Trp Gly Asp His Cys Glu Ile Tyr Pro Cys Pro
260 900 905 910
262 Val Tyr Ser Ser Ala Glu Phe His Ser Leu Cys Pro Asp Gly Lys Gly

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/763,994

DATE: 07/11/2001
 TIME: 11:05:29

Input Set : A:\X-12239SeqList.app
 Output Set: N:\CRF3\07112001\I763994.raw

263	915	920	925
265	Tyr Thr Gln Asp Asn Asn Ile Val Asn Tyr Gly Ile Pro Ala His Arg		
266	930	935	940
268	Asp Ile Asp Glu Cys Met Leu Phe Gly Ser Glu Ile Cys Lys Glu Gly		
269	945	950	955
271	Lys Cys Val Asn Thr Gln Pro Gly Tyr Glu Cys Tyr Cys Lys Gln Gly		960
272	965	970	975
274	Phe Tyr Tyr Asp Gly Asn Leu Leu Glu Cys Val Asp Val Asp Glu Cys		
275	980	985	990
277	Leu Asp Glu Ser Asn Cys Arg Asn Gly Val Cys Glu Asn Thr Arg Gly		
278	995	1000	1005
280	Gly Tyr Arg Cys Ala Cys Thr Pro Pro Ala Glu Tyr Ser Pro Ala Gln		
281	1010	1015	1020
283	Arg Gln Cys Leu Ser Pro Glu Glu Met Glu Arg Ala Pro Glu Arg Arg		
284	1025	1030	1035
286	Asp Val Cys Trp Ser Gln Arg Gly Glu Asp Gly Met Cys Ala Gly Pro		1040
287	1045	1050	1055
289	Leu Ala Gly Pro Ala Leu Thr Phe Asp Asp Cys Cys Cys Arg Gln Gly		
290	1060	1065	1070
292	Arg Gly Trp Gly Ala Gln Cys Arg Pro Cys Pro Pro Arg Gly Ala Gly		
293	1075	1080	1085
295	Ser His Cys Pro Thr Ser Gln Ser Glu Ser Asn Ser Phe Trp Asp Thr		
296	1090	1095	1100
298	Ser Pro Leu Leu Leu Gly Lys Pro Pro Arg Asp Glu Asp Ser Ser Glu		
299	1105	1110	1115
301	Glu Asp Ser Asp Glu Cys Arg Cys Val Ser Gly Arg Cys Val Pro Arg		1120
302	1125	1130	1135
304	Pro Gly Gly Ala Val Cys Glu Cys Pro Gly Gly Phe Gln Leu Asp Ala		
305	1140	1145	1150
307	Ser Arg Ala Arg Cys Val Asp Ile Asp Glu Cys Arg Glu Leu Asn Gln		
308	1155	1160	1165
310	Arg Gly Leu Leu Cys Lys Ser Glu Arg Cys Val Asn Thr Ser Gly Ser		
311	1170	1175	1180
313	Phe Arg Cys Val Cys Lys Ala Gly Phe Ala Arg Ser Arg Pro His Gly		
314	1185	1190	1195
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317	1205		
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321	<211> LENGTH: 3771		
322	<212> TYPE: DNA		
323	<213> ORGANISM: Homo sapiens		
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327	gtgatctgca agcgacactg tctcaaggcc cagtgtcgaa acagttgtca gcagggtctcc 120		
328	aacatgacgc tcatacgaga gaacggccac agcacagaca cgctcacggg ctccggcttc 180		
329	cgcgtgtgg tgtgccctct cccctgcatt aatggccggcc agtgctcctc gcaaaccag 240		
330	tgcctgtgtc ccccgactt cactggccgc ttctgcccagg tgcccccaagg aggagccggt 300		
331	gggggttaccg gcggctcagg ccccgccctg agcaggacag gggccctgtc cacagggcgc 360		
332	tgcggccccc tggctccgga gggcgactt gtggccagca agcacgcatt ctacggcgtc 420		

09/163,444 D

<210> 6
<211> 1257
<212> PRT
<213> Homo sapiens

<220>
<223> Xaa = any amino acid encoding codon or nonsense codon

<400> 6

Xaa cannot represent a nonsense codon -

it can only represent an actual amino acid

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/763,994

DATE 07/11/2001

TIME 11:05:30

Input Set : A:\X-12239SeqList.app
Output Set: N:\CRF3\07112001\I763994.raw

L:10 M:270 C: Current Application Number differs, Replaced Application Number
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:496 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:6
L:496 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:6
L:496 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6